1

## SEQUENCE LISTING

<110>	TAMURA, TOMOHIRO			
<120>	NOVEL EXPRESSION VECTOR SUITABLE FOR PROTEIN AT LOW TEMPERATURE	R EXPRESSION	OF RECOMBINA	ΓN
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	10/524,193 2005-02-11			
	PCT/JP03/10209 2003-08-11	•		
	JP 2002/235008 2002-08-12			
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His His His His His
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gcgtggacgg cgtcagagaa gggagcggcc atg ggc cac cat cac cat cac cat
                                                               174
                               Met Gly His His His His His
atg gga att cta cgt agc ggc cgc gga tcc aag ctt aga tct cga gga
Met Gly Ile Leu Arg Ser Gly Arg Gly Ser Lys Leu Arg Ser Arg Gly
    10
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tgaactagtc gacccacegg caccegtgag cccctegetg egggtgeegg tgegagggac 282 tgcaacacgc gaaacctgca caaacacacg gaggttggaa tgagcgccac ggacacaccc 342 gataccggcg ccgttccacc ccggttggtg accaccgctg gggcggctga cctgctacgc 402 cgcctcagcg ggactctagt <210> 116 <211> 24 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic coded protein sequence <400> 116 Met Gly His His His His His Met Gly Ile Leu Arg Ser Gly Arg 10 Gly Ser Lys Leu Arg Ser Arg Gly <210> 117 <211> 416 <212> DNA <213> Artificial Sequence <220> <221> CDS <222> (151)..(216) <220> <223> Description of Artificial Sequence: Synthetic nucleotide sequence <400> 117 gtgtacatat cgaggcggc tcccacggcc gcccgggctg agggagccga cggcacgcgg 60 gcgtggacgg cgtcagagaa gggagcggcc atg gga att cta cgt agc ggc cgc Met Gly Ile Leu Arg Ser Gly Arg gga tcc aag ctt aga tct cga gga cat cac cat cac cat cac 216 Gly Ser Lys Leu Arg Ser Arg Gly His His His His His 10 15 tgaactagte gacccacegg caccegtgag ecectegetg egggtgeegg tgegagggae 276 tgcaacacgc gaaacctgca caaacacacg gaggttggaa tgagcgccac ggacacaccc 336 gataccggcg ccgttccacc ccggttggtg accaccgctg gggcggctga cctgctacgc 396

416

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cgcctcagcg ggactctagt
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His His His His His
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gcgtggacgg cgtcagagaa gggagcgcat atg ggc cat cac cat cac cat cac
                               Met Gly His His His His His
gcc atg gga att cta cgt agc ggc cgc gga tcc aag cct aga tct cga
                                                              222
Ala Met Gly Ile Leu Arg Ser Gly Arg Gly Ser Lys Pro Arg Ser Arg
                       15
    10
gga tgaactagtc gacccaccgg cacccgtgag cccctcgctg cgggtgccgg
                                                              275
Gly
25
tgcgagggac tgcaacacgc gaaacctgca caaacacacg gaggttggaa tgagcgccac 335
ggacacaccc gataccggcg ccgttccacc ccggttggtg accaccgctg gggcggctga 395
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cctgctacgc cgcctcagcg ggactctagt
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Arg Gly Ser Lys Pro Arg Ser Arg Gly
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                               Met Gly Ile Leu Arg Ser Gly Arg
                                                              216
gga tcc aag ctt aga tct cga gga cat cac cat cac cat cac
Gly Ser Lys Leu Arg Ser Arg Gly His His His His His His
tgaactagtc gacccaccgg cacccgtgag cccctcgctg cgggtgccgg tgcgagggac 276
tqcaacacgc gaaacctgca caaacacacg gaggttggaa tgagcgccac ggacacaccc 336
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His His His His His
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                                                                   47
   Met Gly Ile Leu Arg Ser Gly Arg Gly Ser Lys Leu Arg Ser Leu
                                                            15
                                                                   81
gag cat cac cat cac cat cac tgaactagtc gac
Glu His His His His His
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His His His His His
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20

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gag cat cac cat cac cat cac tgaactagtc gac
                                                                   82
Glu His His His His His
                 20
<210> 126
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His His His His His
             20
<210> 127
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gggcttgcac ctcacgtcac gtgaggaggc agcgtggacg gcgtcagaga agggagcggc 120
catg
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gcccas	adde dateeegeee dateeedaya aggagatada eedeg	
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